

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) Applicant: University of Texas System

(ii) Title of Invention: METHODS AND COMPOSITIONS  
COMPRISING DNA DAMAGING AGENTS  
AND p53

(iii) Number of Sequences: 4

(iv) Correspondence Address:

(A) Addressee: Arnold, White & Durkee  
(B) Street: P.O. Box 4433  
(C) City: Houston  
(D) State: Texas  
(E) Country: USA  
(F) Zip: 77210

(v) Computer Readable Form:

(A) Medium Type: Floppy Disk  
(B) Computer: IBM PC Compatible  
(C) Operating System: PC-DOS/MS-DOS  
(D) Software: ASCII-DOS

(vi) Current Application Data:

(A) Application Number: Unknown  
(B) Filing Date: Submitted herewith  
(C) Classification: Unknown

(vii) Prior Application Data:

(A) Application Number: 08/145,826  
(B) Filing Date: October 29, 1993  
(C) Classification: Unknown

(A) Application Number: 07/960,543  
(B) Filing Date: October 13, 1992  
(C) Classification: Unknown

(A) Application Number: 07/665,538  
(B) Filing Date: March 6, 1991  
(C) Classification: Unknown

(viii) Attorney/Agent Information:

(A) Name: David L. Parker  
(B) Registration Number: 32,165  
(C) Reference/Docket Number: UTSC:403/PAR

5 (ix) Telecommunication Information:

(A) Telephone: (512) 320-7200  
(B) Telefax: (512) 474-7577

10 (2) INFORMATION FOR SEQ ID NO:1:

(i) Sequence characteristics:

15 (A) Length: 20 base pairs  
(B) Type: Nucleic acid  
(C) Strandedness: Single  
(D) Topology: Linear

(ii) Sequence Description: SEQ ID NO:1:

20 GGCCCCACCC CCTTGGCTTC 20

(2) INFORMATION FOR SEQ ID NO:2:

25 (iii) Sequence characteristics:

(A) Length: 20 base pairs  
(B) Type: Nucleic acid  
(C) Strandedness: Single  
(D) Topology: Linear

(iv) Sequence Description: SEQ ID NO:2:

30 TTGTAACCAT, TATAAGCTGC 20

(2) INFORMATION FOR SEQ ID NO:3:

(v) Sequence characteristics:

40 (A) Length: 20 base pairs  
(B) Type: Nucleic acid  
(C) Strandedness: Single  
(D) Topology: Linear

45 (vi) Sequence Description: SEQ ID NO:3:

TCGTTTCTCA GCAGCTGTTG 20

(2) INFORMATION FOR SEQ ID NO:4:

50 (vii) Sequence characteristics:

- (A) Length: 20 base pairs
- (B) Type: Nucleic acid
- (C) Strandedness: Single
- (D) Topology: Linear

5 (viii) Sequence Description: SEQ ID NO:4:

CATCTGAACT CAAAGCGTGG

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